

Fig. 6A

C.anc.env (subtype C ancestral env. The amino acid sequence is different from Los Alamos Database August 2002)

GC CGCCATGCG CGTGATGGGCATCCTGCGCAACTGCCAGCAGTGGTGAT
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ACCA CCTGTCTGCGCCTCCGACG CCAAGGCC TA CGAGCGCGAGGTGCA
CAACGTGTGGGCCACC CA CGCCTGCGTGCC CACCGACCCCAA CCC CCAGG
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ATGGTGGAC CAGATGCACGAGGACATCA TC TCCCTGTGGGAC CAGTCCT
GAAGCCCTGCGTGAAGCTGACCC CCCTGTGCGTGACCTGAACTGCA CCA
ACGTGACCAACGCCACCAACAACACTA CAACGGCGAGATGAAGAAGTGC
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CATCGGCCAGGCCCAC TGCAACATCTCCGAGGCAAGTGGAA CAAGACCC
TGCAGCAGGTGGCCGAGAAGCTGGGCAAGCACTTCCCAA CAAGACCATC
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TCCCTGCTGGA CACCATCGCCATCGCCGTGGCCGAGGGCA CCGACCGCAT
CATCGAGGTGGTGAGCGCGCCTGCCGCGCATCTGAAACATCCCCCGCC
GCATCCGCCAGGGCTTCGAGGCCGCCCTGCTGTAA

Fig. 6B

C.con.env (subtype C consensus env. The amino acid sequence is different from Los Alamos Database August 2002)

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GCCGCCATGCGCGTGATGGGCATCCTGCGCAACTGCCAGCAGTGGTGGAT
CTGGGGCATCCTGGGCTTCTGGATGCTGATGATCTGCAACGTGGTGGGCA
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ACGTGACCAACGCGCACCAACAACACCTACAACGAGGAGATCAAG AACTGC
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CCTGTTCTACCGCCTGGACATCGTGCCCCCTGAACGAGAACTCCTCCGAGT
ACCGCCTGATCAACTGCAACACCTCCGCCATCACCCAGGCCTGCCCAAG
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CATCCTGAAGTGCAACAACAAGACCTTCAACGGCACCCGGCCCCCTG CAACA
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CCGCTCCTCCCTGCGCGGCTGCGAGCGCGGCTGGGAGGCCCTGAAGTACC
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Fig. 8

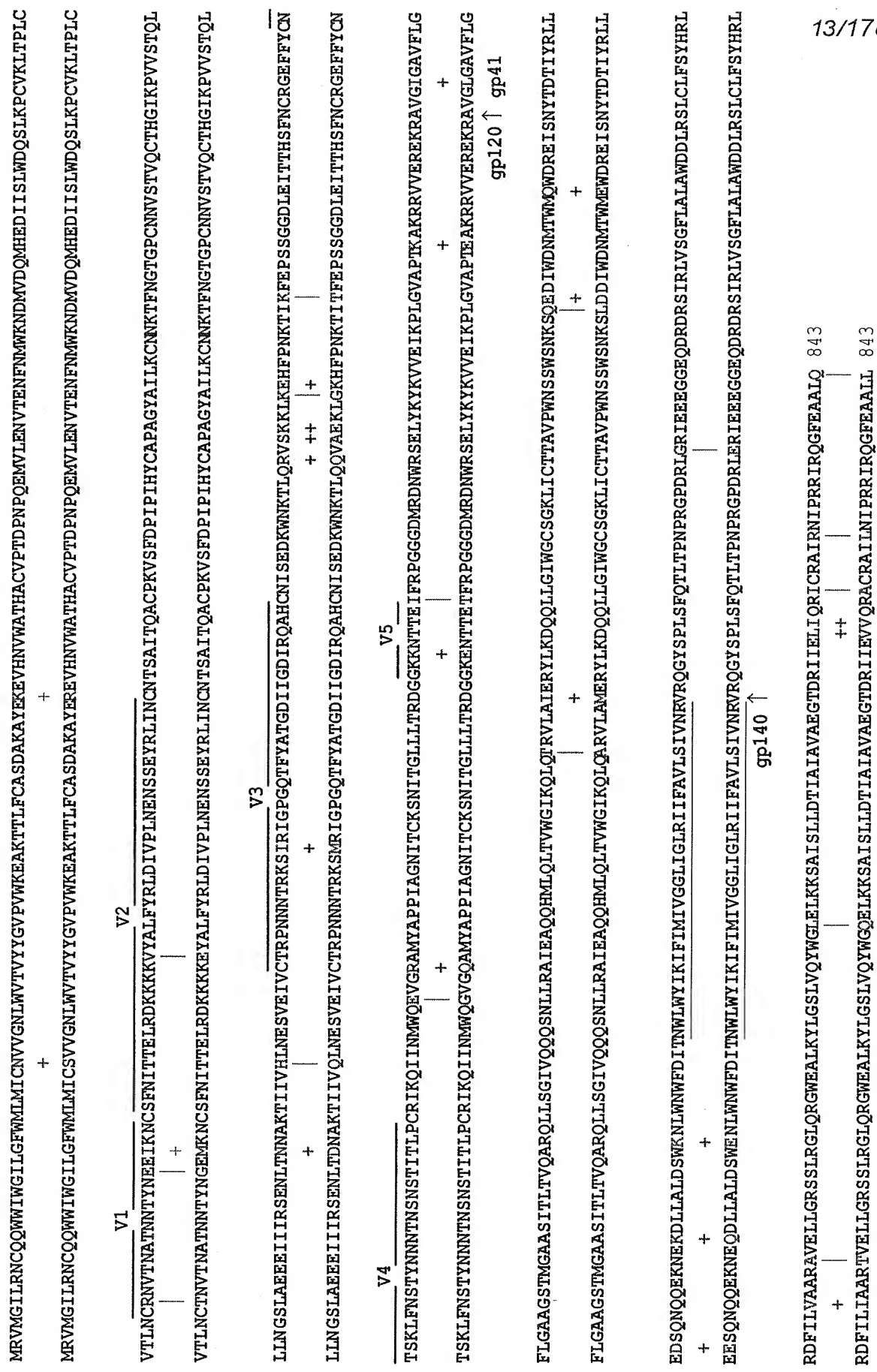
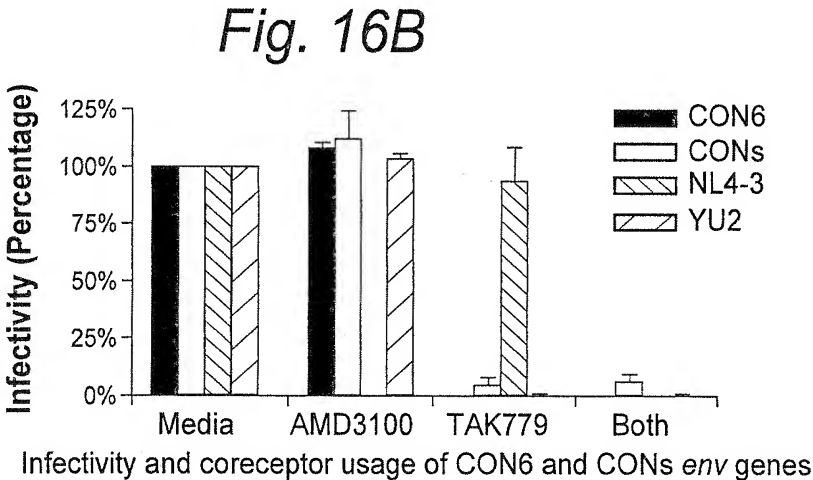
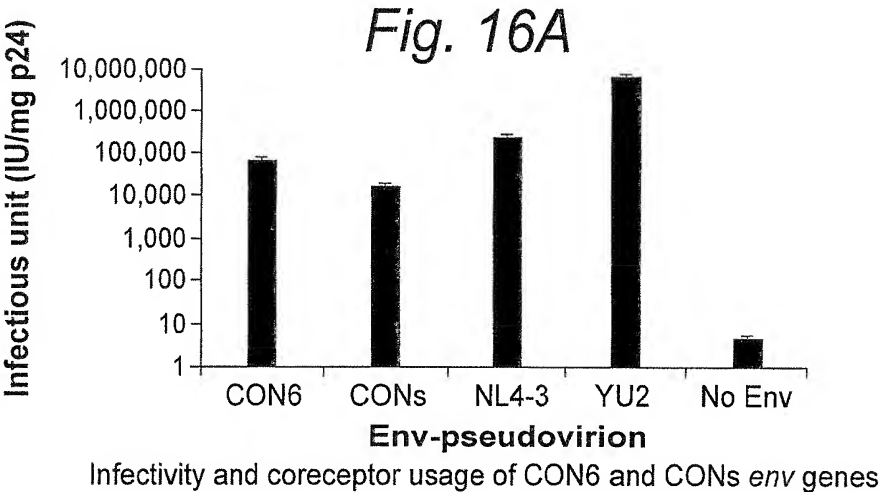
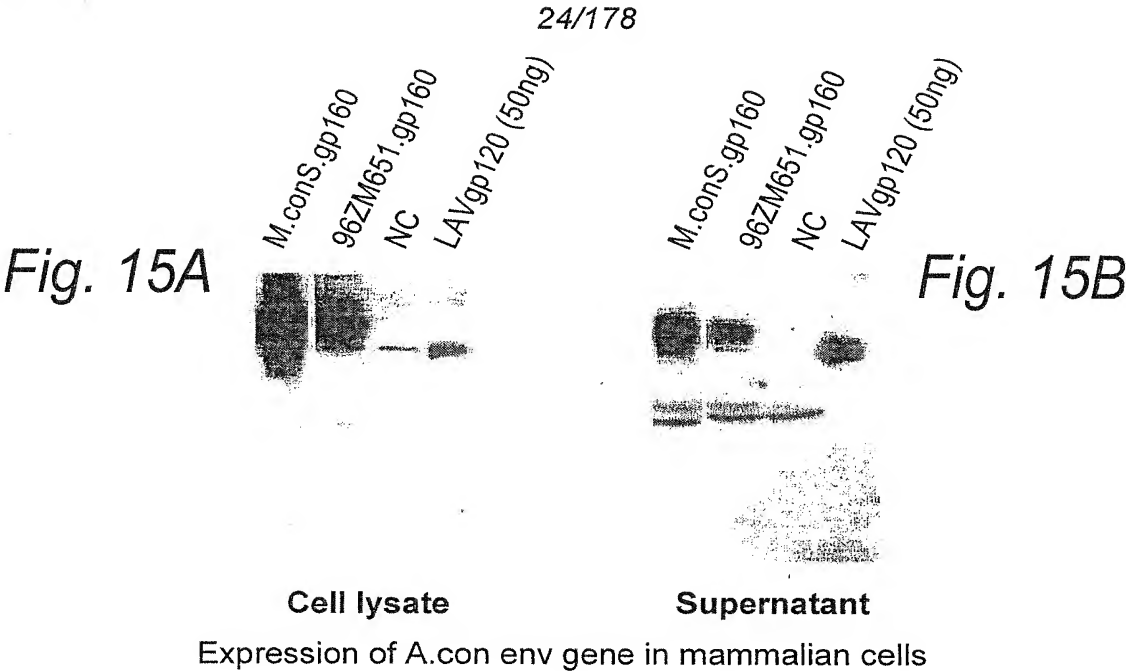


Fig. 14B

CONs.env (gorup M consensus env gene. This one contain the consensus sequence for variable regions in env gene. The identical amino acid sequences as in the public domain)

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GCGCTGGGGCACCCTGATCCTGGGCATGCTGATGATCTGCTCCGCCGCCG
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AACAACACCCGCAAGTCCATCCGCATCGCCCCCGGCCAGGCCTTCTACGC
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GCCTGCGCCGCGGCTGGGAGGCCCTGAAGTACCTGTGGAACCTGCTGCAG
TACTGGGGCCAGGAGCTGAAGAACTCCGCCATCTCCCTGCTGGACACCAC
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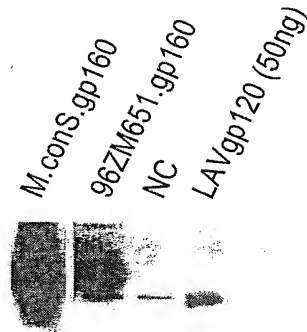
Fig. 18B

A.con.env (subtype A consensus env. Identical amino acid sequence to that in the public domain)

GCCGCCGCCATGCGCGTGATGGGCATCCAGCGCAACTGCCAGCACCTGTG
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GCACAACGTGTGGGCCACCCACGCCTGCGTGCCACCGACCCCAACCCCC
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AACATGGTGGAGCAGATGCACACCGACATCATCTCCCTGTGGGACCAGTC
CCTGAAGCCCTGCGTGGAAGCTGACCCCCCTGTGCGTGACCCCTGAACTGCT
CCAACGTGAACGTGACCACCAACATCACCAACATCACCGACAACATGAAG
GGCGAGATCAAGAACTGCTCCTTCAACATGACCACCGAGCTGCGCGACAA
GAAGCAGAAGGTGTACTCCCTGTTCTACAAGCTGGACGTGGTGCAGATCA
ACAAGTCCAACCTCCTCCTCCAGTACCGCCTGATCAACTGCAACACCTCC
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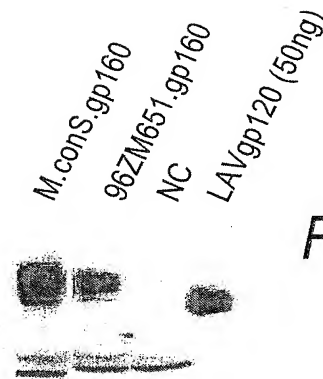
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Fig. 18C



Cell lysate

Fig. 18D



Supernatant

Expression of A.con env gene in mammalian cells

Fig. 19A

M.con.gag (group M consensus gag. Identical amino acid sequence to that in the public domain)

```

GCCGCCGCCATGGGCGCCCGCGCCTCCGTGCTGTCCGGCGGCAAGCTGGA
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TAA
  
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M.con.pol.nuc

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Fig. 19B

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Fig. 19C

M.con.nef (group M consensus nef. Identical amino acid sequence to that in the public domain)

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CGCCGTGCGCGAGCGCATCCGCCGCACCA CCCCGCCGCCGAGGGCTGG
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CTGGAGGGCTGATCTACTCCAAGAAGCGC CAGGAGATCCTGGACCTGTG
GGTGTA CCA CACCCAGGGCTACTTC CCGACTGGCAGAAC TACACCCCG
GCCC CGGCATCGCTA CCCCCTGACCTT CGGCTGGTGCTT CAAGCTGGTG
CCCGTGGACCCCGAGGAGGTGGAGGAGGCCAACGAGGGCGAGAACAACTC
CCTGCTGCA C CCACTGTGCCAGCACGGCATGGAGGACGAGGAGCGCGAGG
TGCTGATGTGGAAGTTCGACTCCCGCTGGCCCTGCGCCACATCGCCCGC
GAGCTGCACC CCGAGTACTACAAGGACTGCTAA

Fig. 19D

C.con.pol.nuc

GCCGCCGCCATGCCCCAGATCACCTGTGGCAGCGCCCCCTGGTGTCCAT
CAAGGTGGGCGGCCAGATCAAGGAGGCCCTGCTGGCCACCGGCGCCGACG
ACACCGTGCTGGAGGAGATCAACCTGCCCGGCAAGTGGAAGCCCAAGATG
ATCGGCGGCATCGGCGGCTTCATCAAGGTGCGCCAGTACGACCAGATCCT
GATCGAGATCTGCGGCAAGAAGGCATCGGCACCGTGCTGGTGGGCCCCA
CCCCCGTGAACATCATCGGCCGCAACATGCTGACCCAGCTGGGCTGCACC
CTGAACCTCCCCATCTCCCCCATCGAGACCGTGCCCGTGAAGCTGAAGCC
CGGCATGGACGGCCCCAAGGTGAAGCAGTGGCCCCCTGACCGAGGAGAAGA
TCAAGGCCCTGACCGCCATCTGCGAGGAGATGGAGAAGGAGGGCAAGATC
ACCAAGATCGGCCCCGAGAACCCCTACAACACCCCCGTGTTCCGCATCAA
GAAGAAGGACTCCACCAAGTGGCGCAAGCTGGTGGACTTCCGCGAGCTGA
ACAAGCGCACCCAGGACTTCTGGGAGGTGCAGCTGGGCATCCCCACCCC
GCCGGCCTGAAGAAGAAGAAGTCCGTGACCGTGCTGGACGTGGGCGACGC
CTACTTCTCCGTGCCCCCTGGACGAGGGCTTCCGCAAGTACACCGCCTTCA
CCATCCCCCTCCATCAACAACGAGACCCCCGGCATCCGCTACCAGTACAAC
GTGCTGCCCCAGGGCTGGAAGGGCTCCCCCGCCATCTTCCAGTCTCTCCAT
GACCAAGATCCTGGAGCCCTTCCGCGCCCGAGAACCCCGAGATCGTGATCT
ACCAGTACATGGACGACCTGTACGTGGGCTCCGACCTGGAGATCGGCCAG
CACCGCGCCAAGATCGAGGAGCTGCGCGAGCACCTGCTGAAGTGGGGCTT
CACCACCCCCGACAAGAAGCACCAAGGAGCCCCCTTCTGTGGATGG
GCTACGAGCTGCACCCCGACAAGTGGACCGTGCAGCCCATCCAGCTGCCC
GAGAAGGACTCCTGGACCGTGAACGACATCCAGAAGCTGGTGGGCAAGCT
GAACTGGGCCTCCAGATCTACCCCGGCATCAAGGTGCGCCAGCTGTGCA
AGCTGCTGCGCGGCGCCAAGGCCCTGACCGACATCGTGCCCCCTGACCGAG
GAGGCCGAGCTGGAGCTGGCCGAGAACCGCGAGATCCTGAAGGAGCCCGT
GCACGGCGTGTA CTACGACCCCTCCAAGGACCTGATCGCCGAGATCCAGA
AGCAGGGCCACGACCAGTGGACCTACCAGATCTACCAGGAGCCCTTCAAG
AACCTCAAGACCGGCAAGTACGCCAAGATGCGCACCGCCCAACCAACGA
CGTGAAGCAGCTGACCGAGGCCGTGCAGAAGATCGCCATGGAGTCCATCG
TGATCTGGGGCAAGACCCCCCAAGTTCCGCCTGCCCATCCAGAAGGAGACC
TGGGAGACCTGGTGGACCGACTACTGGCAGGCCACCTGGATTCCCGAGTG
GGAGTTCTGTGAACACCCCCCCCCCTGGTGAAGCTGTGGTACCAGCTGGAGA
AGGAGCCCATCGCCGGCGCGAGACCTTCTACGTGGACGGCGCCGCCAAC

CGGAGACCAAGATCGGCAAGGCCGGCTACGTGACCGACCGCGGCCGCCA
GAAGATCGTGTCCCTGACCCGAGACCACCAACAGAAACCGAGCTGCAGG
CCATCCAGCTGGCCCTGCAGGACTCCGGCTCCGAGGTGAACATCGTGACC
GACTCCAGTACGCCCTGGGCATCATCCAGGCCAGCCCGACAAGTCCGA
GTCCGAGCTGGTGAACACAGATCATCGAGCAGCTGATCAAGAAGGAGCGCG
TGTAACCTGTCTGGGTGCCGCCCAAGGGCATCGCGGCAACGAGCAG
GTGGACAAGCTGGTGTCTCCGGCATCCGCAAGTGCTGTTCTTGGACGG
CATCGACAAGGCCAGGAGGACGAGAAAGTACCACTCCAACCTGGCGCG
CCATGGCTCCGAGTTCAACCTGCCCCCATCGTGGCCAAGGAGATCGTG
GCCTCTGCGACAAGTGCAGCTGAAGGGGAGGCCATGCACGGCCAGGT
GGACTGCTCCCCGGCATCTGGCAGCTGGAATGCACCCACCTGGAGGGCA
AGATCATCTGTGGTGGCCGTGACCTGGCTCCGGCTACATCGAGGCCGAG
GTGATCCCCCGGAGACCGGCCAGGAGACCGCTACTTTCATCCTGAAGCT
GGCGGCCGTGGCCCGTGAAGGTGATCCACACCGACCAAGGCTCCAAC
TCACCTCCGCCCGCTGAAGCCGCTGCTGGTGGCCGGCATCCAGCAG
GAGTTCGGCATCCCCCTACAACCCCGAGTCCAGGGCGTGTGGAGTCCAT
GAACAAAGAGCTGAAGAAGATCATCGGCCAGTGCAGGCCAGGCCGAGC
ACCTCAAGACCGCGGTGCAGATGGCCGTGTTCATCCACAACCTCAAGCGC
AAGGGCGCATCGCGGTACTCCGCCGGGAGCGCATCATCGACATCAT
CGCCACCGACATCCAGACCAAGGAGCTGCAGAAAGCAGATCATCAAGATCC
AGAACTTCCGCGTGTAACCGGACTCCCGGACCCCACTGGAAGGGC
CCCCCAAGCTGTGTGAAGGGGAGGGCGCGCTGGTGTATCCAGGACAA
CTCCGACATCAAGGTGTGCCCCCGCCGAAGGCCAAGATCATCAAGGACT
ACGGCAAGCAGATGGCCGGGCGGACTGCGTGGCCCGCCCGCAGGACGAG
GACTAA

Fig. 19D (continued)

M.con.gag (group M consensus gag)

MGARASVLSGGKLDWEKIRLRPGGKKYRLKHLVWASRELERFALNPGLLETSEG CKQIIGQLQPA
LQTGSEELRSLYNTVATLYCVHQRIEVKDTKEALEKIEEEQNKSQQTQAAADKGNSSKVSQNYPIVQN
LQGQMVHQAI SPRTLNAWKVIEEKAFSPEVIPMFSALSEGATPQDINTMLNTVGGHQAAQMMLKDTINE
EAAEWDRLHPVHAGPI PPGOMREPRGSDIAGTSTLQEQIAWMTSNPPI PVGEIYKRWI ILGLNKIVRM
SPVSILDIRQGPKEFRDYVDRFFKTLRAEQATQDVKNWMTDTLLVQANPDCKTILKALGPGATLEEMM
TACQGVGGPGHKARVLAHAMSQVTNAAIMQRGNFKGQRRI IKFNCGKEGHIARNCRAPRKKGWCCKG
EGHQMKDCTERQANFLGKIWPSNKGPRGNFLQSRPEPTAPPAESFGFGEIITPSPKQEPKDEPPLTSLK
SLFGNDPLSQ

Fig. 19E